

Sequence Listing

SEQ ID NO. 1

5 KGF (FGF-7)

Genbank Accession No. P21781 *Homo sapiens*

1 m h k w i l t w i l p t l l y r s c f h i i c l v g t i s l a c n d m t p e q m a t n v n c s s p e r h t r s y d y m e  
 10 61 g g d i r v r r l f c r t q w y l r i d k r g k v k g t q e m k n s y n i m e i r t v a g i v a i k g v e s e f y l a  
 121 m n k e g k y a k k e c n e d c n f k e l i l e n h y n t y a s a k w t h n g g e m f v a l n q k g i p v r g k k t k  
 181 k e q k t a h f l p m a i t

SEQ ID NO. 2

15 KGF (FGF-7)

Genbank Accession No. NP\_032034 and I48610 *Mus musculus*

20 1 m r k w i l t r i l p t l l y r s c f h l v c l v g t i s l a c n d m s p e q t a t s v n c s s p e r h t r s y d y m e  
 61 g g d i r v r r l f c r t q w y l r i d k r g k v k g t q e m k n s y n i m e i r t v a g i v a i k g v e s e y y l a  
 121 m n k e g k y a k k e c n e d c n f k e l i l e n h y n t y a s a k w t h s g g e m f v a l n q k g i p v k g k k t k  
 181 k e q k t a h f l p m a i t

25 SEQ ID NO. 3

KGF-2 (FGF-10)

30 Genbank Accession No. BAA22331 *Homo sapiens*

1 m w k w i l t h c a s a f p h l p g c c c c f l l l f l v s s p v t c q a l g q d m v s p e a t n s s s s s f s s p  
 61 s s a g r h v r s y n h l q g d v r w r k l f s f t k y f l k i e k n g k v s g t k k e n c p y s i l e i t s v e i g  
 121 v a v k a i n s n y y l a m n k k g k l y g s k e f n d c k l k e r i e e n g y n t y a s f n w q h n g r q m y v a l  
 35 181 n g k g a p r r g q k t r k n t s a h f l p m v v h s

SEQ ID NO. 4

40 KGF-2 (FGF-10)

Genbank Accession No. AAH48229 *Mus musculus*

1 m w k w i l t h c a s a f p h l p g c c c c f l l l f l v s s p v t c q a l g q d m v s q e a t n c s s s s s f s s  
 45 61 p s s a g r h v r s y n h l q g d v r w r r l f s f t k y f l t i e k n g k v s g t k n e d c p y s v l e i t s v e i g  
 121 v v a v k a i n s n y y l a m n k k g k l y g s k e f n d c k l k e r i e e n g y n t y a s f n w q h n g r q m y v a  
 181 l n g k g a p r r g q k t r k n t s a h f l p m t i q t

50 SEQ ID NO. 5

N-terminal Glp-Leu-Gly-Pro-Gln-Gly-Pro-Pro-His-Leu-Val-Ala-Asp-Pro-Ser-Lys-Lys-Gln-Gly-Pro-Trp-  
 Leu-Glu-Glu-Glu-Glu-Glu-Ala-Tyr-Gly-Trp-Met-Asp-Phe

55 SEQ ID NO. 6

N-terminal Glp-Leu-Gly-Pro-Gln-Gly-Pro-Pro-His-Leu-Val-Ala-Asp-Pro-Ser-Lys-Lys-Gln-Gly-Pro-Trp-  
 Leu-Glu-Glu-Glu-Glu-Glu-Ala-Tyr-Gly-Trp-Leu-Asp-Phe

SEQ ID NO. 7

N-terminal Glp-Gly-Pro-Trp-Leu-Glu-Glu-Glu-Glu-Glu-Ala-Tyr-Gly-Trp-Met-Asp-Phe .

5 SEQ ID NO. 8

N-terminal Glp-Gly-Pro-Trp-Leu-Glu-Glu-Glu-Glu-Glu-Ala-Tyr-Gly-Trp-Leu-Asp-Phe

SEQ ID NO. 9

10

mqrhcyyvli falalaafse aswkprsqpp daplggtanr dlelpwleqq gpashhrrql  
gpqgpphlva dpskkqgpwl eeeeeaygwm dfgrrsaede n

SEQ ID NO. 10

15

MCNDMTPEQMATNVNCS SPERHTRS YDMEGGDIRVRRLFCRTQWYLRIKRGKVKGTQEMKNN  
YNIMEIRTVAVGIVAIGVSEFYLAMNKEGKLYAKKECNEDCNFKELILENHYNTYASAKWTHNG  
GEMFVALNQKGI PVRGKKTKEQKTAHFLPMAIT

20 SEQ ID NO. 11

MLGQDMVSPE ATNSSSSSFS SPSSAGRHRV SYNHLQGDVR WRKLFSFTKY  
FLKIEKNGKV SGTKKENCYPY SILEITSVEI GVVAVKAINS NYYLAMNKKG  
KLYGSKEFNN DCKLKERIEE NGYNTYASFN WQHNGRQMYV ALNGKGAPRR  
25 GQKTRRKNTS AHFLPMVVHS